Figure 1A.

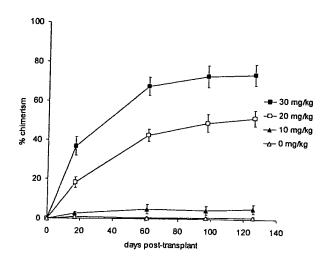


Figure 1B.

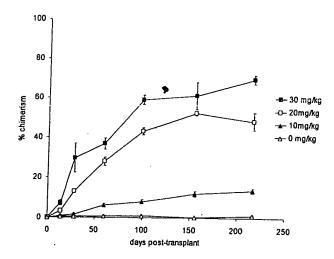


Figure 1C.

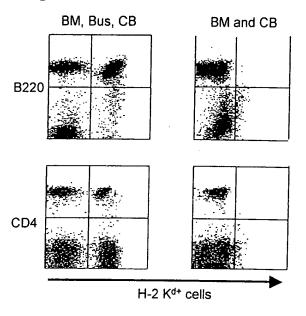
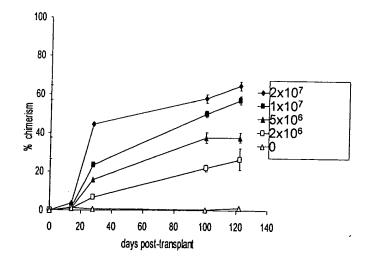


Figure 1D.





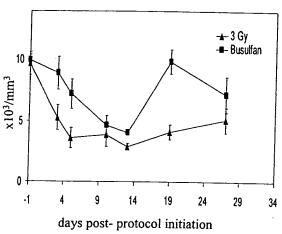


Figure 3A.

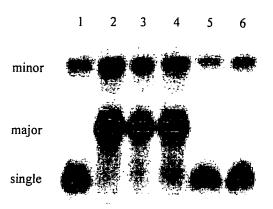


Figure 3B.

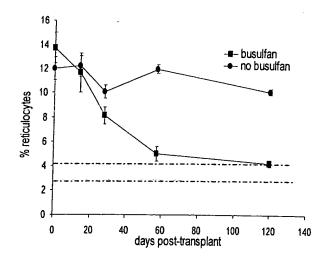


Figure 4A.

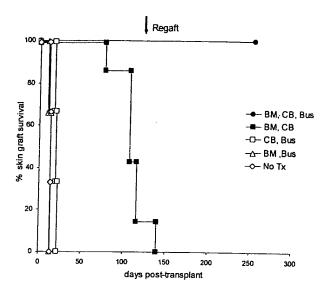


Figure 4B.

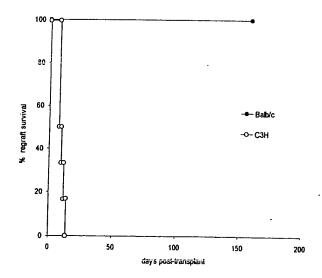


Figure 5A.

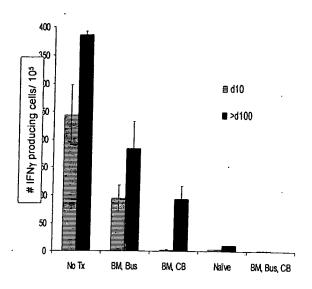


Figure 5B.

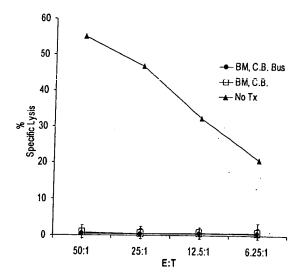


Figure 5C.

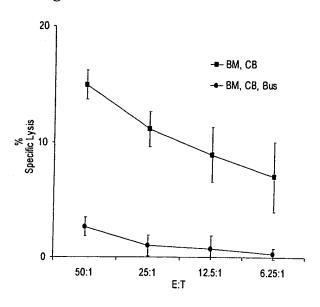


Figure 5D.

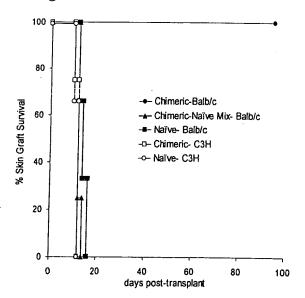


Figure 6A.

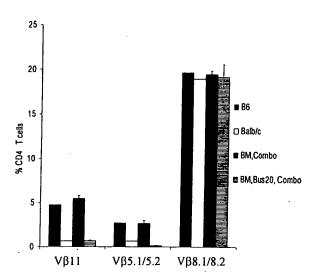


Figure 6B.

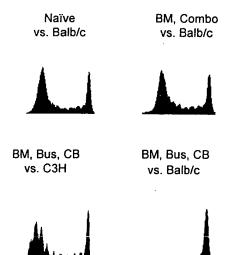
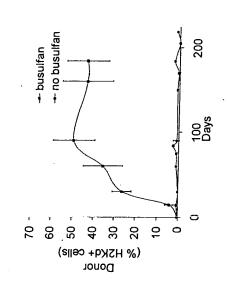


Figure 7a





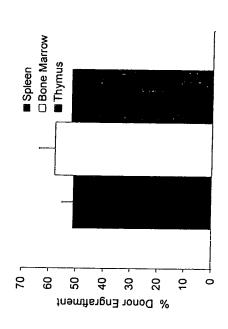


Figure 8

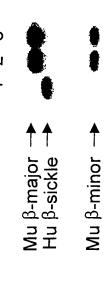
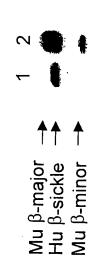
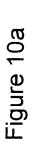
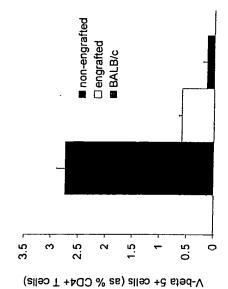


Figure 9









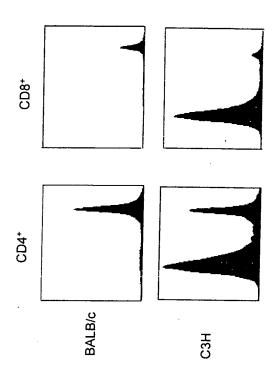
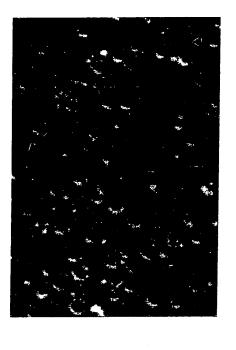


Figure 11

a.





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Figure 11c

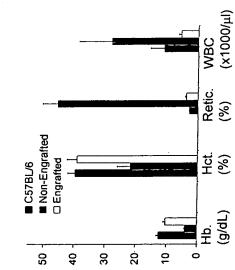
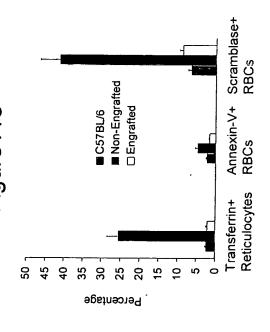
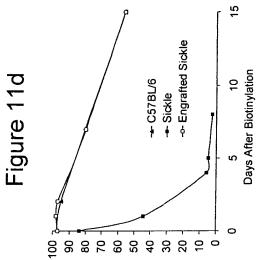


Figure 11e





% of Peripheral Blood Cells Biotinylated

Figure 12a

■ C57BU6 ■ Sickle □ Engrafted Sickle Splenic WBCs Splenic Retics. Splenic RBCs 5 0 70 8 9 20 6 30 20 Percent

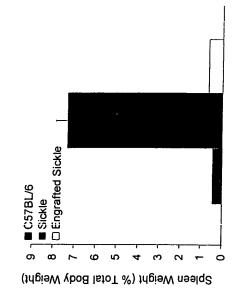
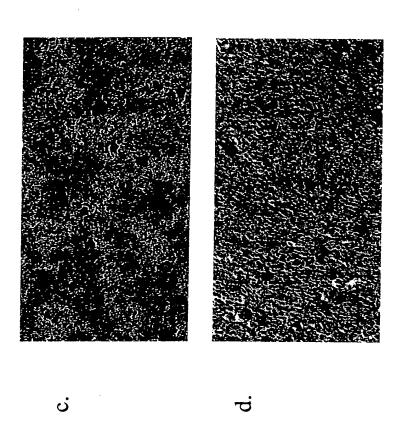
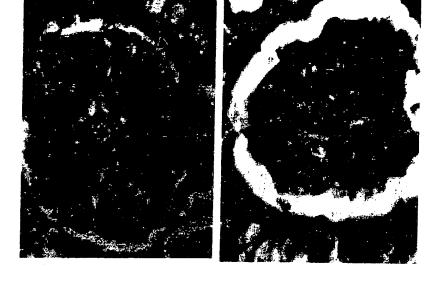


Figure 12b

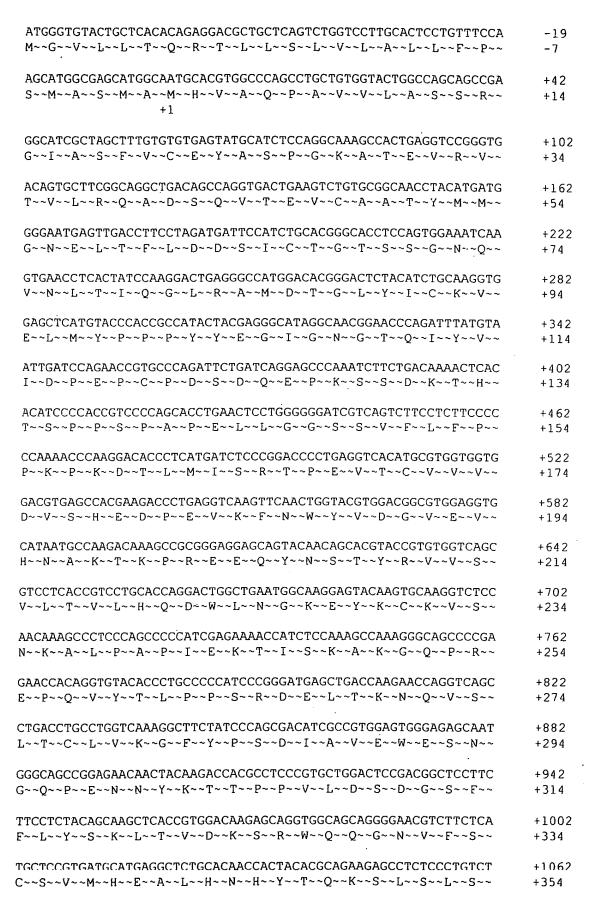
Figure 12



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þ.



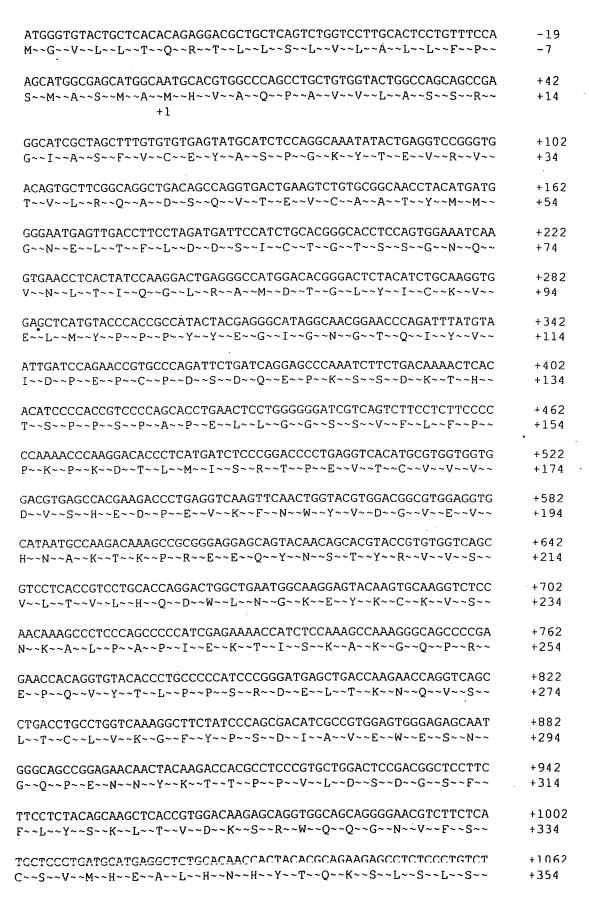


FIG. 16

AGCA S~~M	TGG(CGA	GCA'	TGG	CAAT "Y +i	rgca Th	. ~ V.	rgg A	ccc. ~~Q	AGC ~~p	CTG ~~A	CTC V	TGG V	TAC	TGG	CCAG	CAG	CCGA
GGCA G~~I																		
ACAG T~V																		
GGGA. G~~N																		
GTGA V~~N																		
gagc' e~~l																		
ATTG																		
ACATY T~S																		
CCAAJ P~~K	AACC	CAA ~K~	GGA D~	CAC TT	CCT L	CAT	TAD ~I~	CTC S	CCC	GA(occ.	CTG ~~e	AGG ~~v	TCA(CAT(CGT V	GOTC	GTG V
GACGT D~~V																		
CATA																		
GTCCT V~~L~	CAC TT	CGT ~V~	CCT ~L~	GCA THT	CCA	GGA(TG:	GCT L	GAA "K"	TGC ~G~	CA.	red red	AGT/	ACA/	\GTC	CAA	OGTC	TCC
AACAA N~~K~	AGC	CCT L	CCC ~p~	AGC A	CCC	CATO	GA(GAA K	AAC ~T~	CAT	CTC S	CA	AAG(CAA	\AGG	GCA Q~	GCCC - p	CGA R
GAACC E P																		
CTGAC L~~T~	CTG	CTY	V	CAA/ ~K~	AGGC	F	TA?	rcc P	CAG ~S~	CGA D	CAT	CGC	CGI	NGGA	GTG "W"	GGA	SAGC	AAT Y
GGGCA G~~Q~																		
TTCCT F~~L~	CTAC	AGC	AAC K~	CTC L	TACC	GTG V~~	GAC D~-	AAC K"	GAG(R	gtg "W"	GCA Q	GCA ~Q~	.GGG	gaa 'n'	CGIC	TTC	TCA S
rgctc(
ccccci																		-
						_		_		_								

ATGGGTGTACTGCTCACACAGAGGACGCTGCTCAGTCTGGTCCTTGCACTCCTGTTTCCA M-G-V-L-T-Q-R-T-L-L-S-L-V-L-A-L-L-F-P AGCATGGCGAGCATGCACGTGGCCCAGCCTGCTGTGGTACTGGCCAGCAGCCGA S--N--Y--R--W--H--A--G--B--Y--A--A--Z--Z--Z--Z--OGCATCGCTAGCTTTGTGTGTGAGTATGCATCTCCAGGCAAAACTACTGAGGTCCGGGTG -II-A-S-F-V-C-E-Y-A-S-P-G-X-T-T-E-V-R-V-ACAGTGCTTCGGCAGGCTGACAGCCAGGTGACTGAAGTCTGTGCGGCAACCTACATGATG TTTVTLTTRTQTAATDTTSTQTTVTTTTETTVTTCTAATATTTTYTTMTYT. GGGAATGAGTTGACCTTCCTAGATGATTCCATCTGCACGGGCACCTCCAGTGGAAATCAA GT-NT-ET-LT-TH-FT-LT-DT-DT-ST-IT-CT-TT-GT-TT-ST-ST-GT-NT-O-GTGAACCTCACTATCCAAGGACTGAGGGCCATGGACACGGGACTCTACATCTGCAAGGTG V~~N~~L~~I~Q~~G~~L~~R~~A~~M~~D~~T~~G~~L~~Y~~I~~C~~K~~V~ GAGCTCATGTACCCACCGCCATACTACGAGGGCATAGGCAACGGAACCCAGATTTATGTA ATTGATCCAGAACCGTGCCCAGATTCTGATCAGGAGCCCAAATCTTCTGACAAAACTCAC ITTOTTPTTBTTPTTCTTPTTDTTSTTDTTQTTBTTPTTKTTSTTSTTDTTKTTTTHT ACATCCCCACCGTCCCCAGCACCTGAACTCCTGGGGGGATCGTCAGTCTTCCTCTTCCCC T--S--P--P--S--P--A--P--E--L--L--G--G--S--S--V--F--L--F--P-CCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTG PTKTPTKTDTTTLTMTITTSTRTTPTETTVTTTCTTVTVTVT GACGTGAGCCACGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGCGTGGAGGTG DTTVTTSTTHTETTDTTPTTETTVTKTTFTTNTWTTYTTVTTDTTGTTVTTETTVT CATAATGCCAAGACAAAGCCGCGGGAGGAGCAGTACAACAGCACGTACCGTGTGGTCAGC HTTNTATKTTTKTTPTRTETTBTQTYTNTSTTTTYTRTVTVTS-GTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGGTGCAAGGTCTCC VTTLTTTTVTTLTHTQTTDTWTTTTNTGTTKTTETTYTKTTCTKTTVTS AACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGA NT-KT-AT-LT-PT-AT-PT-IT-ET-KT-TT-IT-ST-KT-AT-KT-GT-QT-PT-R GAACCACAGGTGTACACCCTGCCCCCATCCCGGGATGAGCTGACCAAGAACCAGGTCAGC ETPTQTVTTTTLTPTPTGTRTDTBTTTTTKTNTQTVTS CTGACCTGCCTGGTCAAAGGCTTCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAAT LTTTCTTLTVTKTGTFTTYTPTTSTDTTLTTATVTBTWTTTTSTNT GGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTC G-Q-P-E-N-N-Y-K-T-T-P-P-P-V-L-D-S-D-G-S-P-TTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGGAACGTCTTCTCA F--F--S--K--F--S--K--K--Z--K--K--Z--K--M--Q--Q--G--N--V--F--S--TGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCT CTTSTTVTTMTTHTTETTATTLTTHTTMTTHTTYTTTQTTKTTSTTLTTSTTLTTS CCGGGTAAATGA~~~

FIG. 17

ACAGTGCTTCGGCAGGCTGACAGCCAGGTGACTGAAGTCTGTGCGGCAACCTACATGATG T-V-L-R-Q-A-D-S-Q-V-T-E-V-C-A-A-T-T-Y-M-M-GGGAATGAGTTGACCTTCCTAGATGATTCCATCTGCACGGGCACCTCCAGTGGAAATCAA G-N-B-T-T-F-T-D-D-S-T-C-T-G-T-S-S-S-G-N-0-GTGAACCTCACTATCCAAGGACTGAGGGCCATGGACACGGGACTCTACATCTGCAAGGTG V-N-L-T-1-0-G-L-R-A-M-D-T-G-L-Y-1-C-K-V-GAGCTCATGTACCCACCGCCATACTACGAGGGCATAGGCAACGGAACCCAGATTTATGTA ATTGATCCAGAACCGTGCCCAGATTCTGATCAGGAGCCCAAATCTTCTGACAAAACTCAC I--D--B--B--b--C--b--D--2--D--0--B--b--K--2--2---K--4--ACATCCCCACCGTCCCCAGCACCTGAACTCCTGGGGGGATCGTCAGTCTTCCTCTTCCCC T--S--P--P--S--P--A--P--E--L--C--G--G--S--S--V--F--L--P--P-CCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTG PT-K-P-K-D-T-L-M-II-S-R-T-P-E-V-T-C-V-V GACGTGAGCCACGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTG DTTVTSTTHTTETTDTTPTTRTTVTKTTFTTNTTWTTYTTVTTDTTGTTVTTRTTVT CATAATGCCAAGACAAAGCCGCGGGAGGAGCAGTACAACAGCACGTACCGTGTGGTCAGC H--N--A--K--T--K--P--R--E--E--Q--Y--N--S--T--Y--R--V--V--S-GTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGTCTCC V"-L"-T"-V"-L"-H"-Q"-D"-W"-L"-N"-G"-K"-E"-Y"-K"-C"-K"-V"-S AACAAAGCCCTCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCGA N~K~A~L~P~A~P~I~E~K~T~I~S~K~A~K~G~O~P~R~ GAACCACAGGTGTACACCCTGCCCCCATCCCGGGATGAGCTGACCAAGAACCAGGTCAGC CTGACCTGCCTGGTCAAAGGCTTCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAAT L--T--C--L--V--K--G--F--Y--P--S--D--I--A--V--E--W--E--S--N-GGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTC G--Q--P--E--W--W--K--M--T--P--P--V--D--S--D--G--S--F TTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCA TTLTTYTTBTTKTTLTTTTVTTDTTKTTBTTRTWTTQTTQTTQTTYTTPTTST TGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCT CT-8-TV-MT-HT-ET-A-LT-HT-NT-HT-YT-TT-QT-KT-8-L--S-L--S-CCGGGTAAATGA FIG. 18

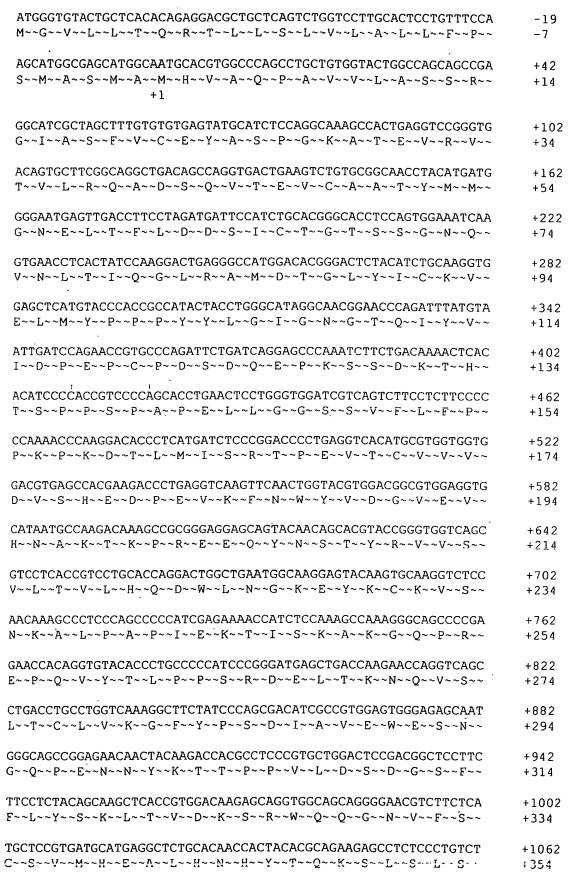
ATGGGTGTACTGCTCACACAGAGGACGCTGCTCAGTCTGGTCCTTGCACTCCTGTTTCCA

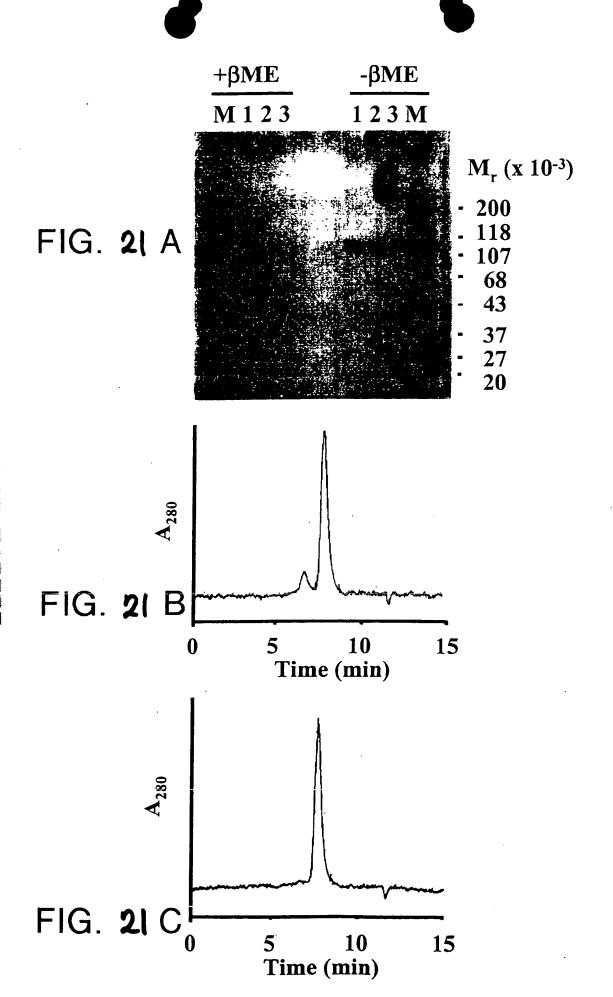
AGCATGGCGAGCATGCCAATGCACGTGGCCCAGCCTGCTGTGGTACTGGCCAGCAGCCGA

GGCATCGCTAGCTTTGTGTGTGAGTATGCATCTCCAGGCAAATGGACTGAGGTCCGGGTG

ONCOSTATIN M SIGNAL PEPTIDE

	G CG GGT								CTG	CTC	AGT				45
A GC	L CA CTC	L CTG	F TTT	P CCA	S AGC	M ATG	A GCG	S AGC	. M	A	+1 M ATG	H CAC	V GTG	A GCC	90
Q CA	P G CCT	A GCT	V GTG	V GTA	L CTG	A GCC	S AGC	S AGC	R CGA	G GGC	I ATC	A GCC	S AGC	F TTT	135
V GT	C G TGT	E GAG	Y TAT	A GCA	s TCT	P CCA	G GGC	K AAA	A GCC	T ACT	E GAG	V GTC	R CGG	V GTG	180
T AC	V A GTG	L CTT	R CGG	Q CAG	A GCT	D GAC	S AGC	Q CAG	V GTG	T ACT	E GAA	V GTC	C TGT	A GCG	225
A GC.	T A ACC	Y TAC	M ATG	M ATG	G GGG	N AAT	E GAG	L TTG	T ACC	F TTC	L CTA	D GAT	D GAT	s TCC	270
I ATG	c TGC	T ACG	G GGC	T ACC	s TCC /	S AGT	G GGA .	и ТAA	Q CAA (V GTG	N AAC	L CTC	T ACT	I ATC	315
Q CAP	G L GGA	L CTG	R . AGG (A I GCC I	M I ATG (D 7	r (ACG (G) GGA (L :	Y : PAC :	ATC :	rgc /	AAG (GTG	360
E GAG	L CTC	M : ATG :	Y I	CA C	o p	CA T	Y YAC I	! I	TG G	GC A		. 5	s (<u>4</u> SITE
T	Q CAG	I \\ TTA	(V PAT G	' I	D TT G	P AT C	E CA G	AA C	CG T	GC C	CA G	S AT T	CT G	AC	450
F TTC	L I	L W	I I	L TC C	A TT G	A CA G	V CA G	S IT A	S GT T	ල ලෙ ලෙ	L GG T	F TG T	P TT T	TT	495
Y TAT	S I	F L	TC C	T C A	A CA GO	V CT G1	s TT TC	L CT T	S IG A	K SC Al	M AA A?	L rg c:	K FA AJ	A G	540
K AAA	R S AGA A	GC C	L CT CT	Т ОА ТЭ	T CA AC	G CA GG	y G GI	Y C TA	V AT GI	K EG AA	M TA A	'G CC	P CC CC	CA	585
T ACA	E P GAG C	E CA G?	C AA TG	E T GA	K A AA	Q G CA	F A TT	Q T CA	P G CC	Y T TA	F T TT	I T AT	P T CC	ec ec	630
I ATC															636





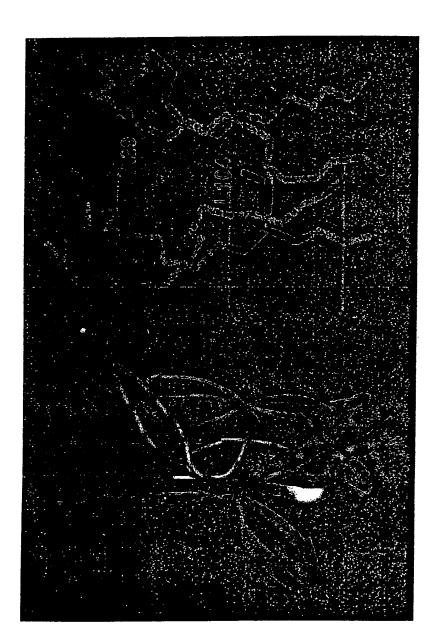
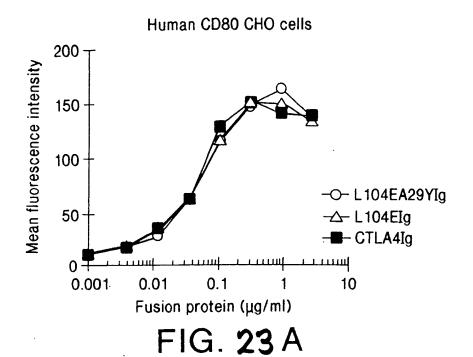
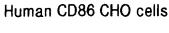
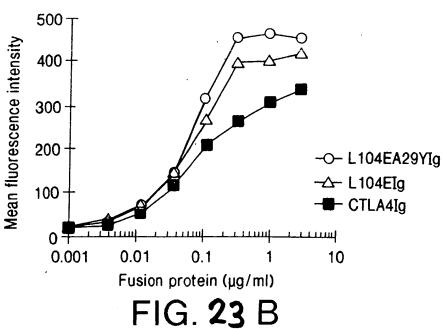
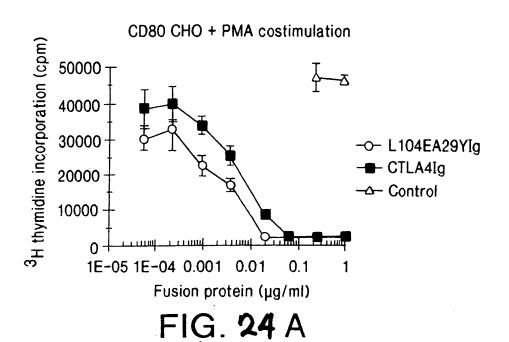


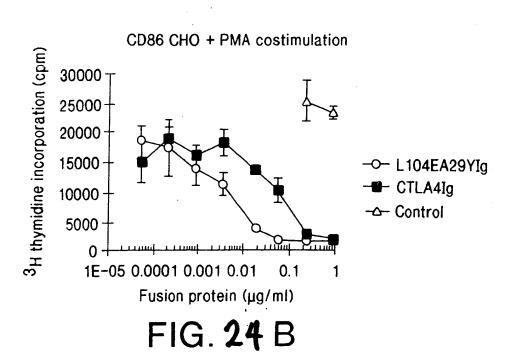
FIG. 22

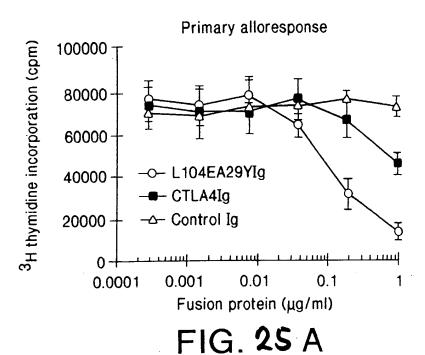


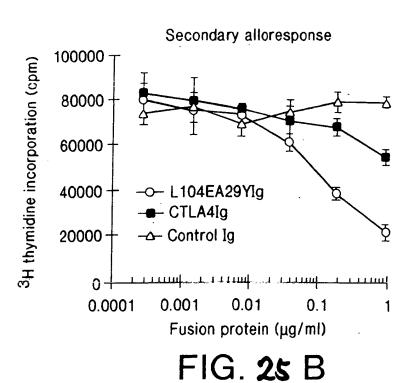












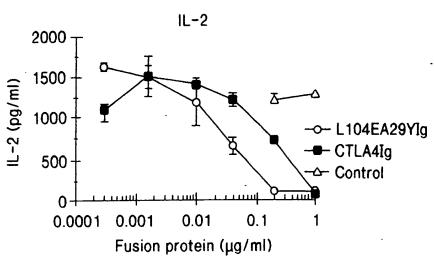


FIG. 26 A

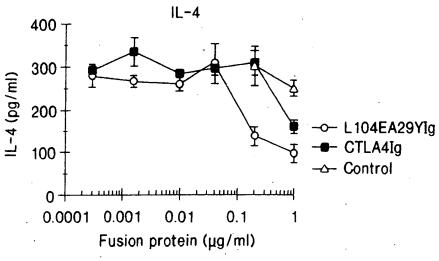


FIG. 26 B

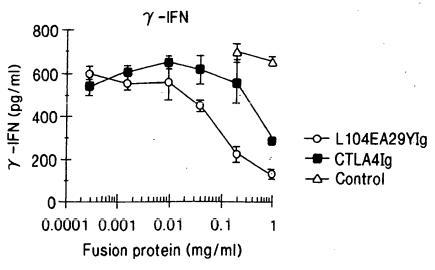
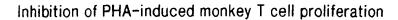


FIG. 26 C



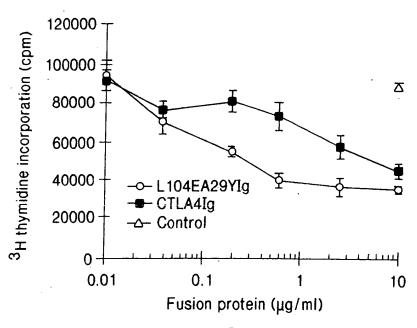
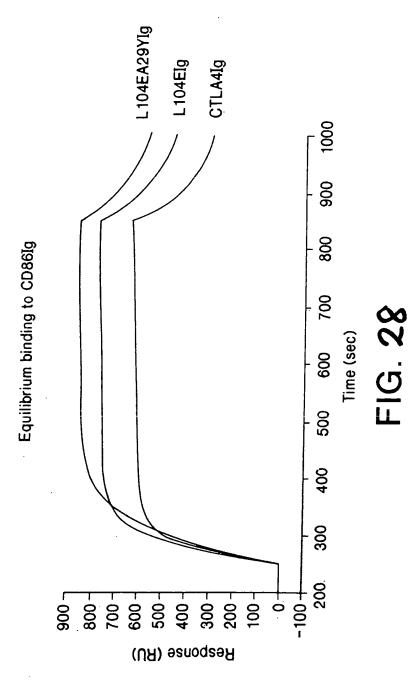


FIG. 27



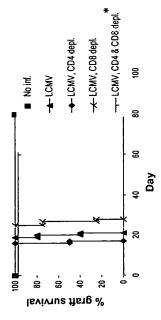


Figure 29

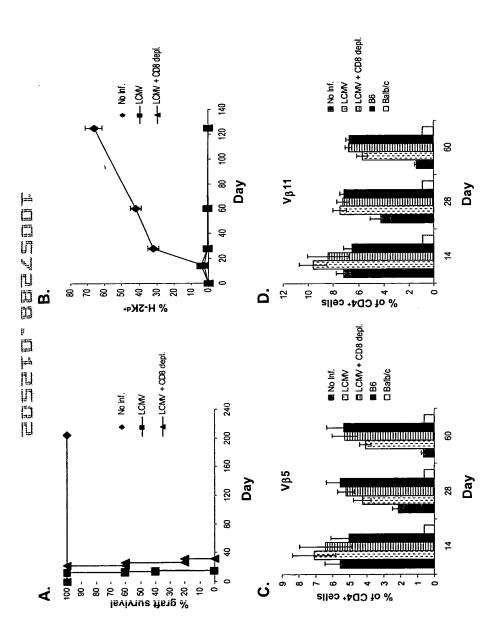


Figure 30

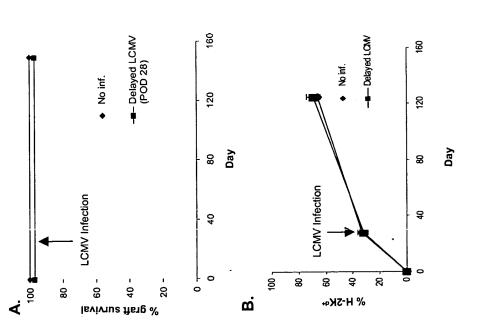


Figure 31

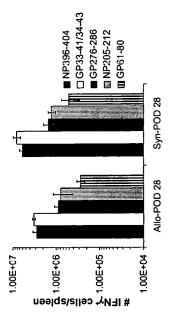


Figure 32

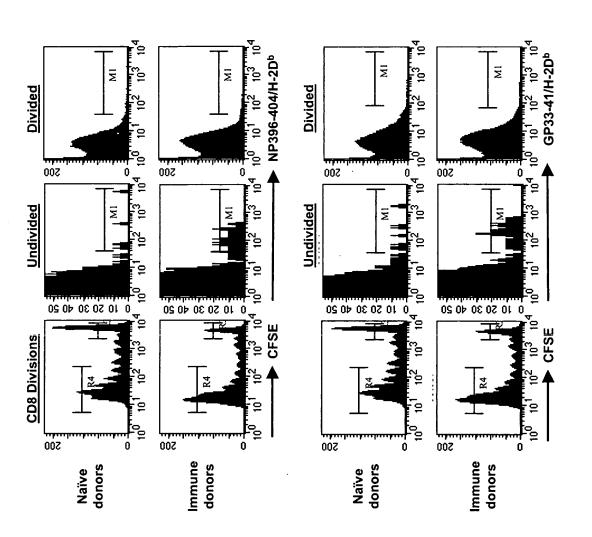


Figure 33

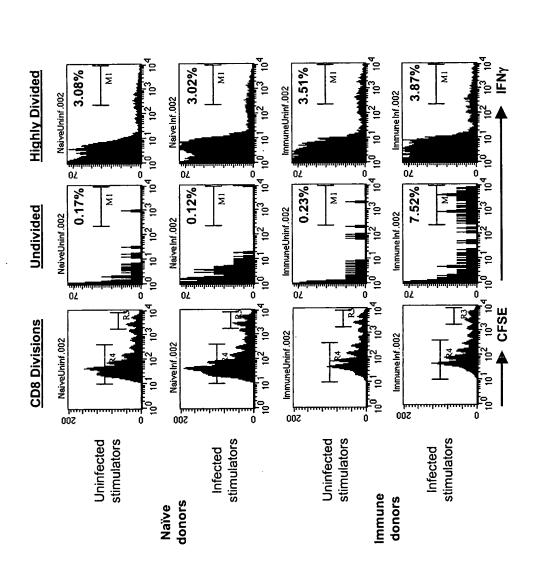


Figure 34



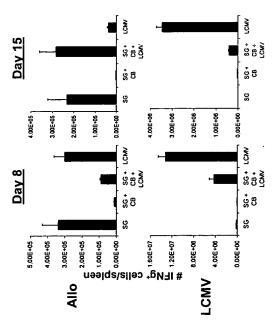


Figure 35

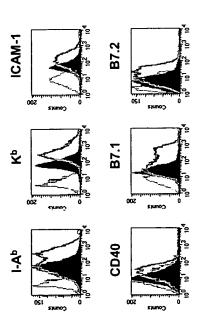


Figure 36